

SEQUENCE LISTING

<110> Chaudhary, Preet M.

<120> Gene Expression in Ectodermal Dysplasia

<130> utsd680

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<170> PatentIn Ver. 2.1

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Gly Asp Cys Arg Gln Gln Glu Phe Arg Asp Arg Ser Gly Asn Cys Val

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Pro Cys Asn Gln Cys Gly Pro Gly Met Glu Leu Ser Lys Glu Cys Gly

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Phe Lys Glu Asp Trp Gly Phe Gln Lys Cys Lys Pro Cys Leu Asp Cys

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		115	120	125	
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	Val Gly Phe Gln Asp Met Glu Cys Val Pro Cys Gly Asp Pro Pro Pro				
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	Pro Tyr Glu Pro His Cys Ala Ser Lys Val Asn Leu Val Lys Ile Ala				
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	Ser Ala Leu Ala Thr Val Leu Leu Ala Leu Leu Ile Leu Cys Val Ile				
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	Tyr Cys Lys Arg Gln Phe Met Glu Lys Lys Pro Ser Trp Ser Leu Arg				
		195	200	205	
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	Ser Gln Asp Ile Gln Tyr Asn Glu Thr Glu Leu Ser Cys Phe Asp Arg				
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	tgt gag gag gcc tgc agc ccc aac ccg gcg act ctt ggt tgt ggg gtg				877
	Cys Glu Glu Ala Cys Ser Pro Asn Pro Ala Thr Leu Gly Cys Gly Val				
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	His Ser Ala Ala Ser Leu Gln Ala Arg Asn Ala Gly Pro Ala Gly Glu				
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320 325 330 335

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340 345 350

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Asn Phe Thr Ala Ala Thr Asp Leu Ser Arg Tyr Asn Asn Thr Leu Val
370 375 380

20 gaa tca gca tca act cag gat gca cta act atg aga agc cag cta gat 1261
Glu Ser Ala Ser Thr Gln Asp Ala Leu Thr Met Arg Ser Gln Leu Asp
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 35 40 45
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 50 55 60
 Gly Tyr Gly Glu Asp Ala Gln Cys Val Ala Cys Arg Leu His Arg Phe
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 Lys Glu Asp Trp Gly Phe Gln Lys Cys Lys Pro Cys Leu Asp Cys Ala
 10 85 90 95
 Val Val Asn Arg Phe Gln Lys Ala Asn Cys Ser Ala Thr Ser Asp Ala
 100 105 110
 Ile Cys Gly Asp Cys Leu Pro Gly Phe Tyr Arg Lys Thr Lys Leu Val
 115 120 125
 15 Gly Phe Gln Asp Met Glu Cys Val Pro Cys Gly Asp Pro Pro Pro Pro
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 Tyr Glu Pro His Cys Ala Ser Lys Val Asn Leu Val Lys Ile Ala Ser
 145 150 155 160
 Thr Ala Ser Ser Pro Arg Asp Thr Ala Leu Ala Ala Val Ile Cys Ser
 20 165 170 175
 Ala Leu Ala Thr Val Leu Leu Ala Leu Leu Ile Leu Cys Val Ile Tyr
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 Cys Lys Arg Gln Phe Met Glu Lys Lys Pro Ser Trp Ser Leu Arg Ser
 195 200 205
 25 Gln Asp Ile Gln Tyr Asn Glu Thr Glu Leu Ser Cys Phe Asp Arg Pro
 210 215 220
 Gln Leu His Glu Tyr Ala His Arg Ala Cys Cys Gln Cys Arg Arg Asp
 225 230 235 240
 Ser Val Gln Thr Cys Gly Pro Val Arg Leu Leu Pro Ser Met Cys Cys
 30 245 250 255
 Glu Glu Ala Cys Ser Pro Asn Pro Ala Thr Leu Gly Cys Gly Val His
 260 265 270
 Ser Ala Ala Ser Leu Gln Ala Arg Asn Ala Gly Pro Ala Gly Glu Met
 275 280 285
 35 Val Pro Thr Phe Phe Gly Ser Leu Thr Gln Ser Ile Cys Gly Glu Phe
 290 295 300
 Ser Asp Ala Trp Pro Leu Met Gln Asn Pro Met Gly Gly Asp Asn Ile
 305 310 315 320

Ser Phe Cys Asp Ser Tyr Pro Glu Leu Thr Gly Glu Asp Ile His Ser
 325 330 335
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 340 345 350
 5 Gln Asp Leu Val Gly Gly Ala Val Pro Val Gln Ser His Ser Glu Asn
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 Phe Thr Ala Ala Thr Asp Leu Ser Arg Tyr Asn Asn Thr Leu Val Glu
 370 375 380
 Ser Ala Ser Thr Gln Asp Ala Leu Thr Met Arg Ser Gln Leu Asp Gln
 10 385 390 395 400
 Glu Ser Gly Ala Val Ile His Pro Ala Thr Gln Thr Ser Leu Gln Val
 405 410 415
 Arg Gln Arg Leu Gly Ser Leu
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